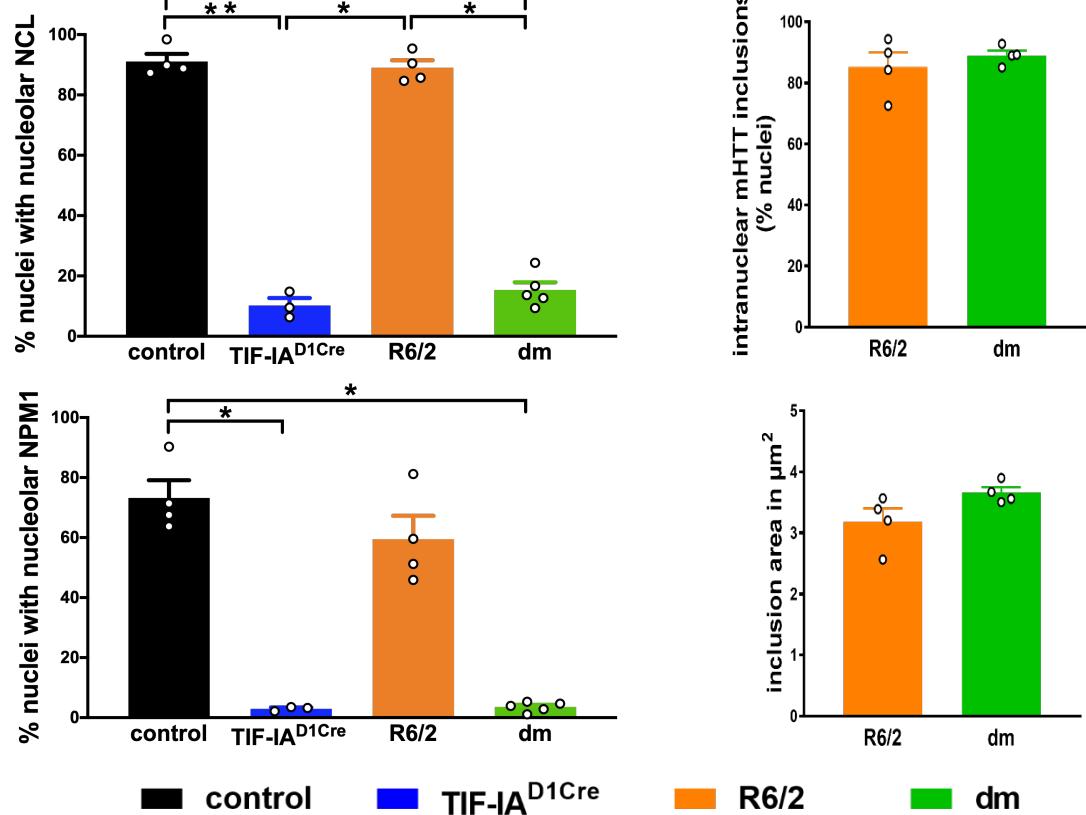
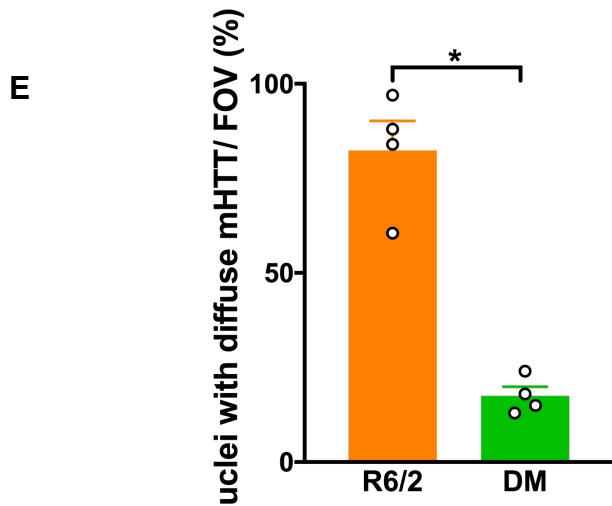
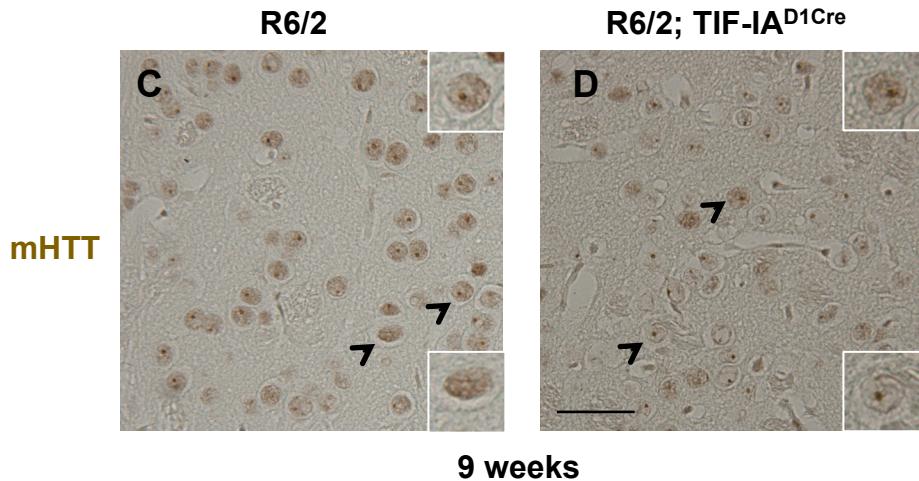
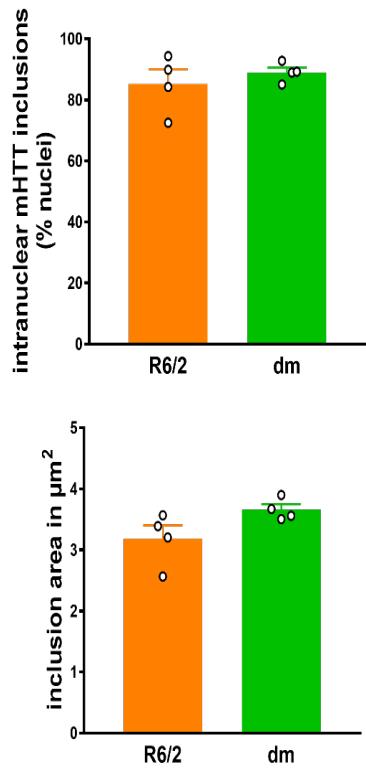
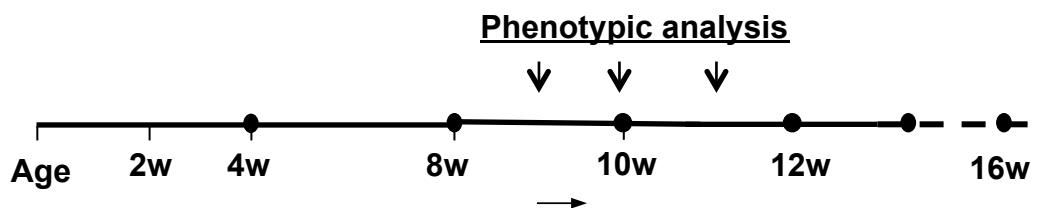
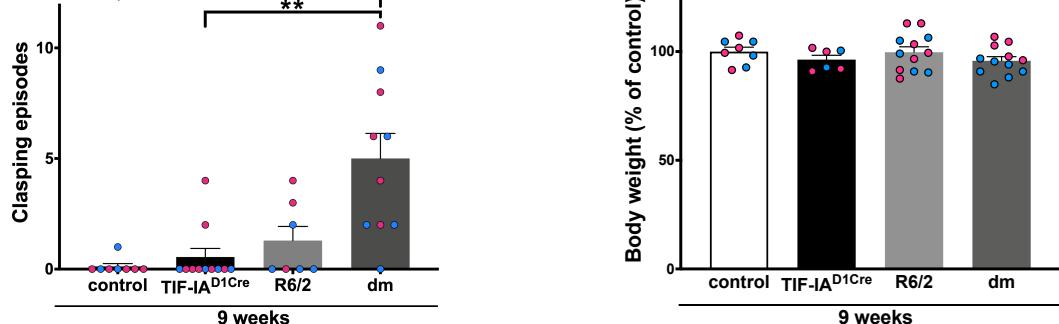
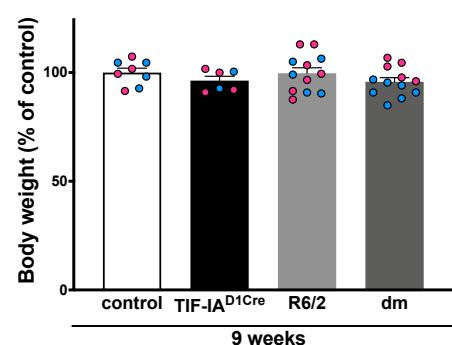
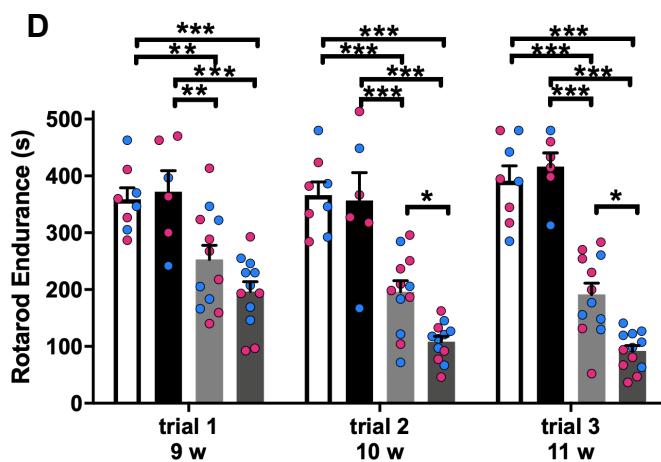
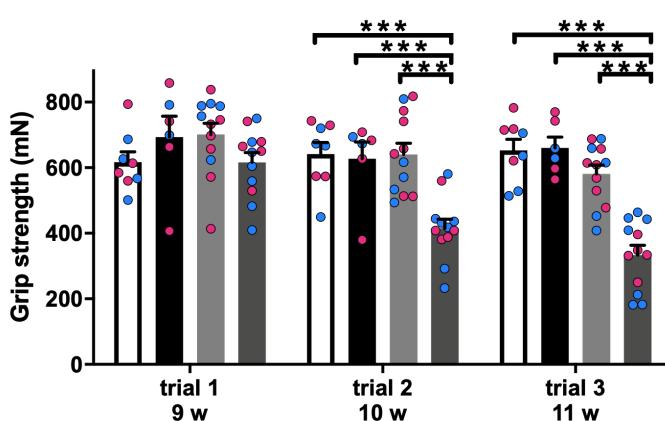


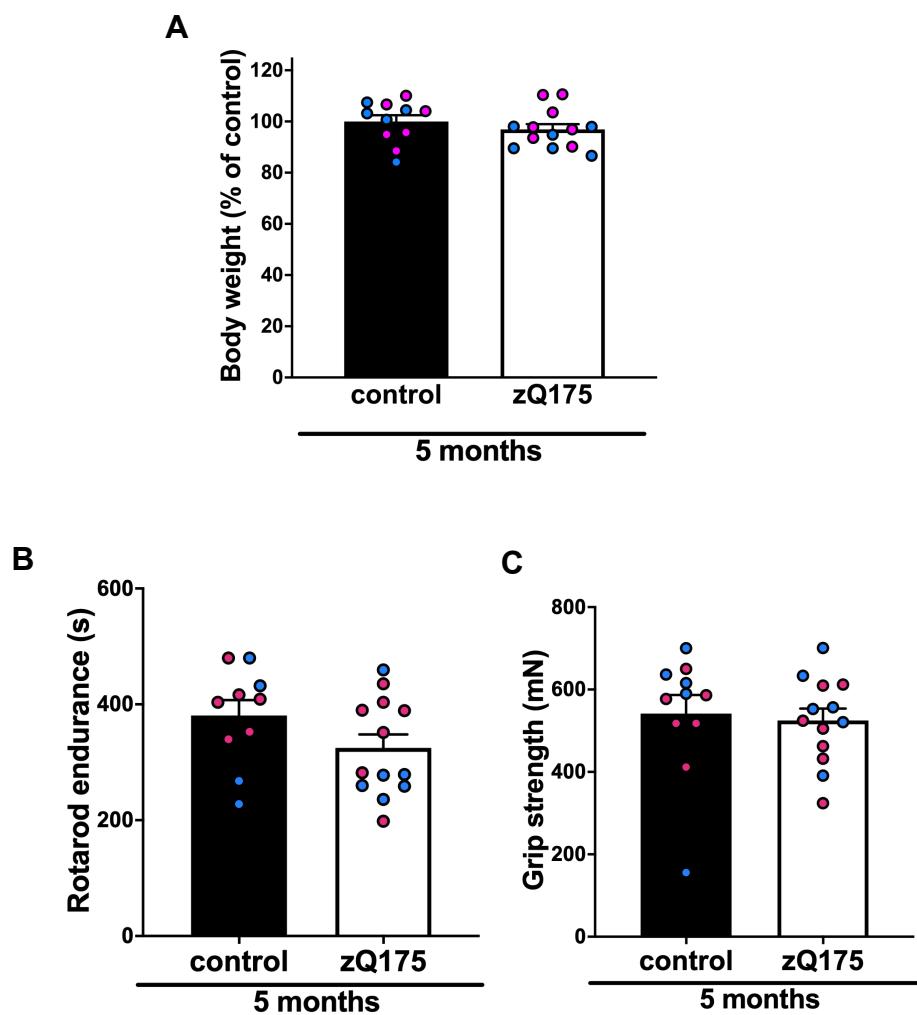
Suppl. Figure 1

A**B****Suppl. Figure 2**

A

□ control ■ TIF-IA^{D1Cre} ■ R6/2 ■ dm

B**C****D****E**

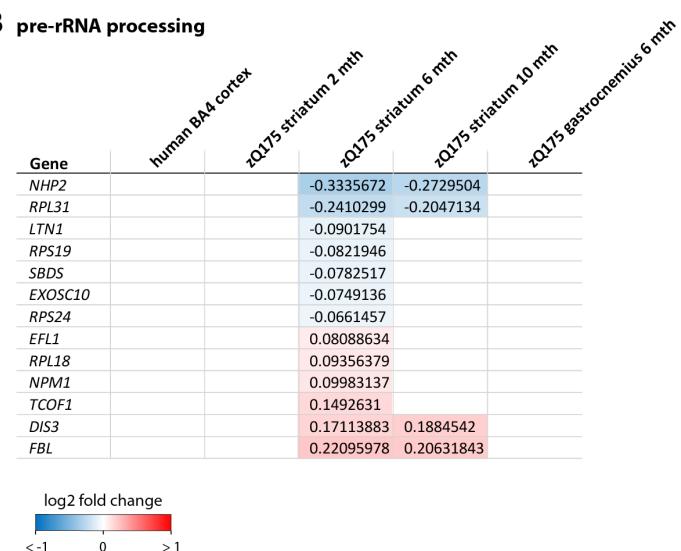


Suppl. Figure 4

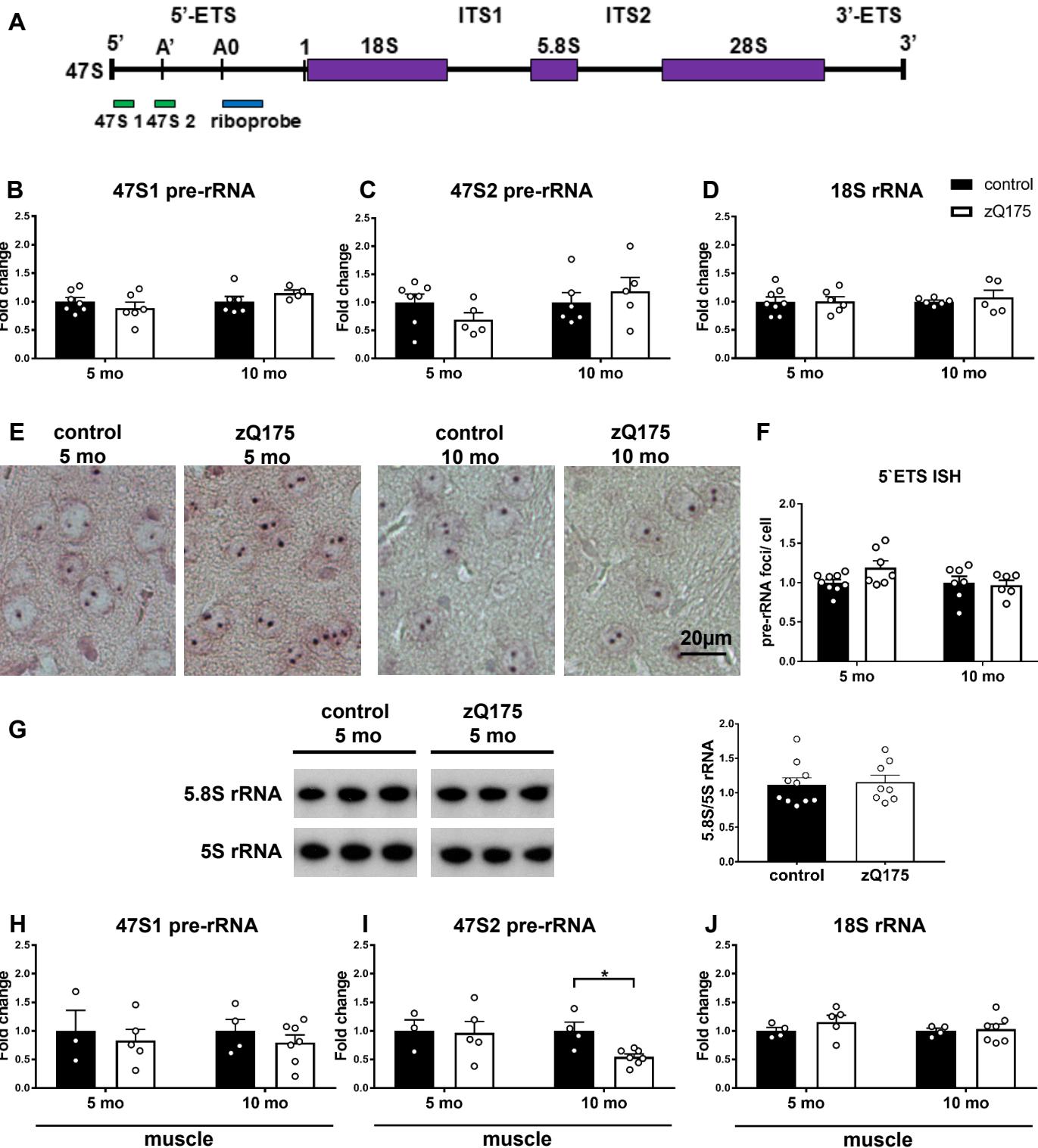
A rRNA transcription

Gene	human BA4 cortex	zQ175 striatum 2 mth	zQ175 striatum 6 mth	zQ175 striatum 10 mth	zQ175 gastrocnemius 6 mth
<i>RRN3</i>	0.29313315				
<i>UBTF</i>	-0.1757289	-0.320152	-0.252921		
<i>GTF2H2</i>			-0.129181		
<i>ERCC6</i>	-0.1514219	-0.1519405			
<i>TAF1C</i>	-0.0894431				
<i>GTF2H3</i>	-0.0754702				
<i>ERCC2</i>	0.06336042				
<i>PCNA</i>	0.09297059				
<i>KAT2B</i>	0.10771751	0.12790748			
<i>GTF2H5</i>	0.10827059				
<i>DDX21</i>	0.11621982	0.18567179			
<i>TOP1</i>	0.11746182				
<i>POLR1A</i>	0.14489715				
<i>ERCC3</i>	0.17364773	0.23119398			
<i>MYO1C</i>	0.18749787	0.2273397			
<i>POLR1E</i>	0.21095236				
<i>XRCC5</i>	0.24006573	0.19054009			

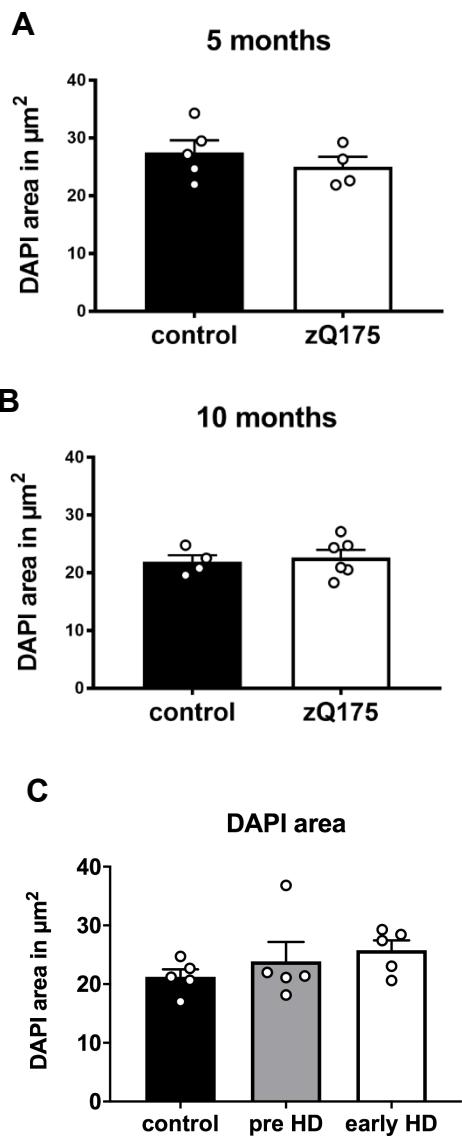
B pre-rRNA processing



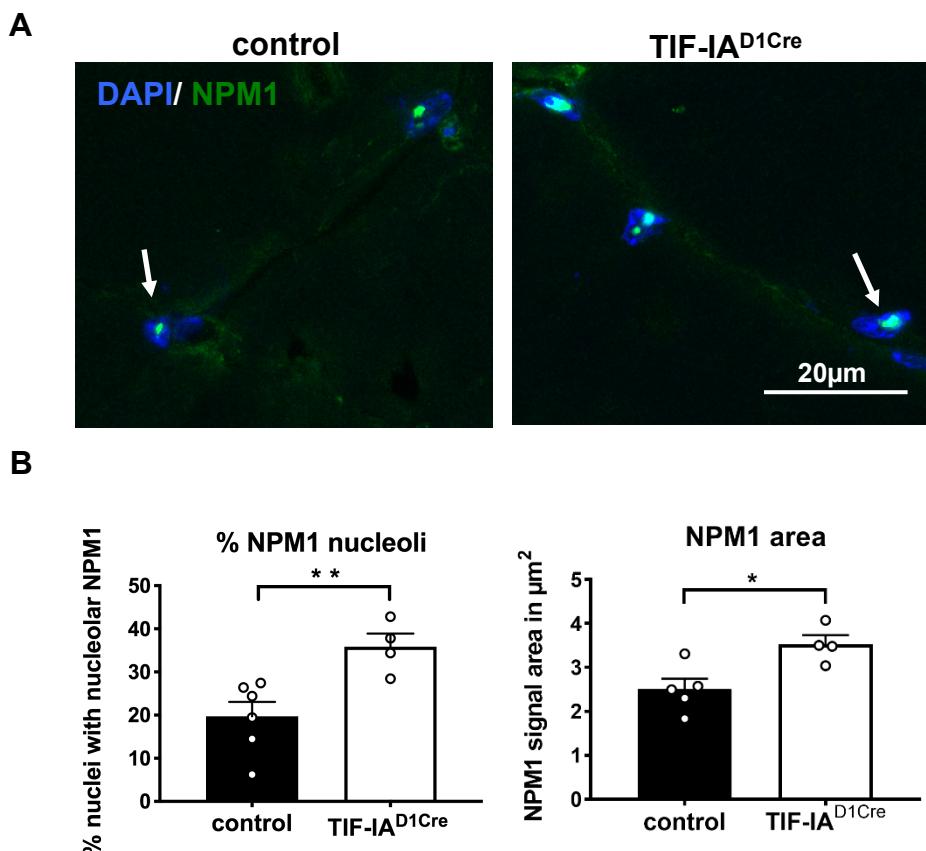
Suppl. Figure 5



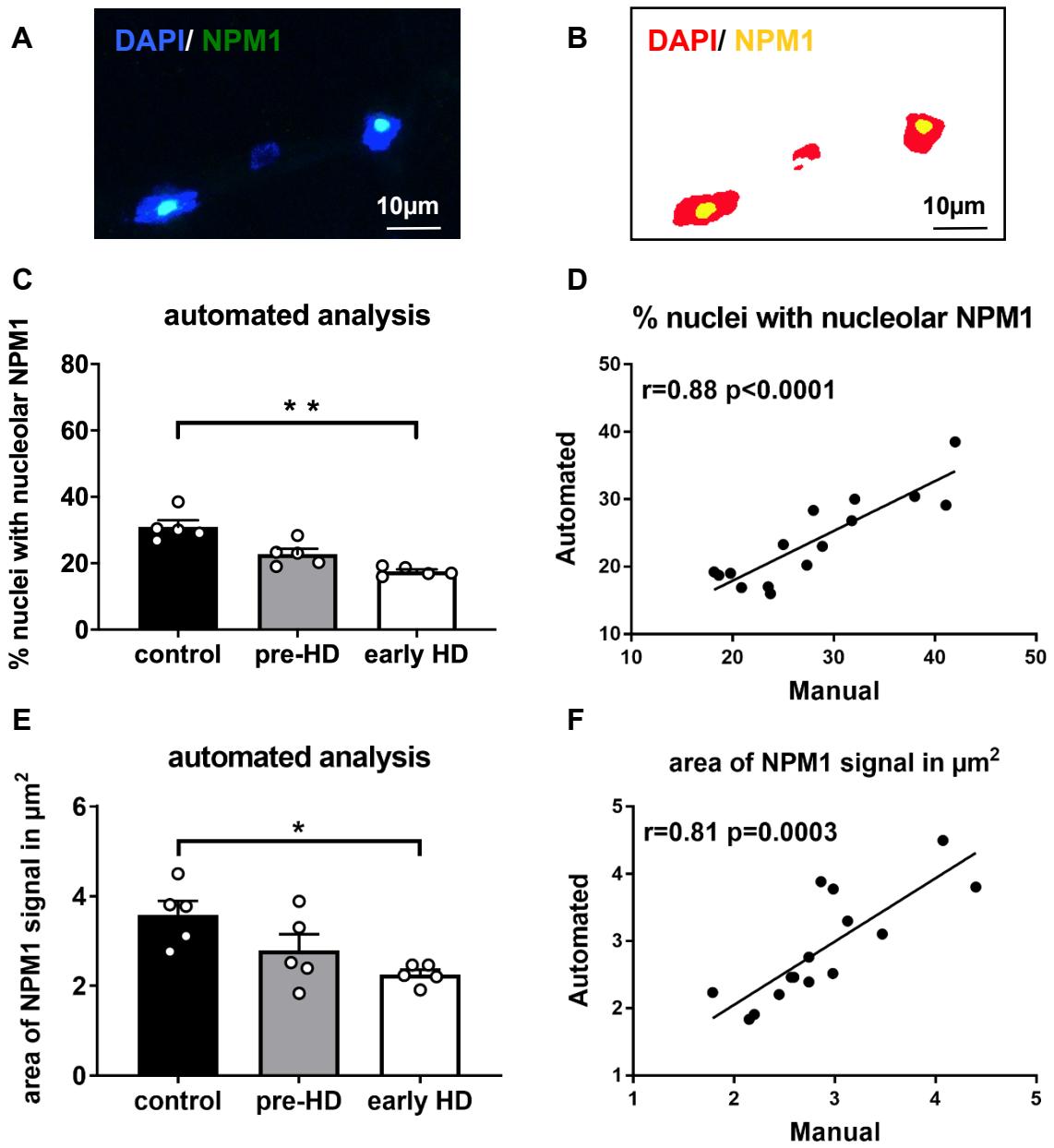
Suppl. Figure 6



Suppl. Figure 7



Suppl. Figure 8



Suppl. Figure 9